## SEQUENCE LISTING

<110> Wang, Tianlin Lohuis, Michael M. Kojima, Cheryl J. Du, Fengxing Byatt, John C. <120> MARKER ASSISTED BEST LINEAR UNBIASED PREDICTION (MA-BLUP): SOFTWARE ADAPTIONS FOR PRACTICAL APPLICATIONS FOR LARGE BREEDING POPULATIONS IN FARM ANIMAL SPECIES <130> 11916.0065.00PC00 <150> US 60/543,034 <151> 2004-02-09 <160> 3 <170> PatentIn version 3.3 <210> 1 <211> 5888 <212> DNA <213> Sus scrofa <220> <221> misc\_feature <222> (1)..(5888) <223> Sequence for AMPK gamma subunit <400> 1 atgagettee tagageaagg agagageegt teatggeeat eeegagetgt aaceaceage 60 tcagaaagaa gccatgggga ccaggggaac aaggcctcta gatggacaag gcaggaggat 120 gtagaggaag gggggcctcc gggcccgagg gaaggtgagt tcaaggccag ttctggggag 180 ctgggactgg gggcagtggg cagtcctcaa acctggggcc cgtctctggt ctggtccctc 240 cataacacag gcacataaca tcatgcagcc agtctcccca caagggggag gacaactgca 300 ttgctgatcc aggggtccag ggatccaagg tggccaactc aggacagagc cactgtcttc 360 tetgtgacte tetgagacte agetetetea cetgcaaaat ggggccacag cattcagget 420 tcccaaggtt gcaatgagga tgaatggaga cagcagatga ggaagttctc tggaagaggg 480 agttactgtg ctctccctcc cgctccccga acaggtcccc agtccaggcc agttgctgag 540 tecaceggge aggaggeeae atteceeaag gecacaceet tggeecaage egeteeettg 600 geegaggtgg acaaeeeeee aacagagegg gacateetee cetetgaetg tgeageetea 660 gcctccgact ccaacacaga ccatctggat ctgggcatag agttctcagc ctcggcggcg 720 tegggggatg agettggget ggtggaagag aageeageee egtgeeeate eecagaggtg 780 ctgttaccca ggctgggctg ggatgatgag ctgcagaagc cgggggccca ggtctacatg 840

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